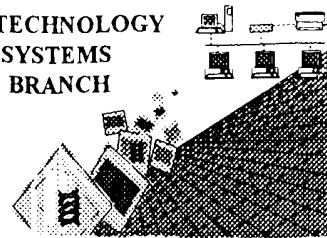


030

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/766,880A
Source: OIF
Date Processed by STIC: 12/13/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

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- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. **EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)**
2. **U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202**
3. **Hand Carry directly to:**
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
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Revised 01/29/2002

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OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/766,880A

DATE: 12/13/2002
TIME: 15:12:56

Input Set : A:\PTO.txt
Output Set: N:\CRF4\12132002\I766880A.raw

W--> 2 <110> APPLICANT:
 W--> 3 Jones, Chris L.
 W--> 4 Abrams, Ezra S.
 W--> 5 Kieffer-Higgins, Stephen G.
 W--> 6 Zhang, Tianhong
 W--> 7 Eacock, Graham P.
 W--> 8 Webb, Michael
 W--> 9 McDowell, Christopher S.
 W--> 14 <110> APPLICANT:
 W--> 15 Methods of and Apparatus for Separating and Detecting Nucleic Acid
 W--> 20 <130> FILE REFERENCE:
 W--> 21 2313.2001-001
 C--> 26 <140> CURRENT APPLICATION NUMBER:
 W--> 27 US 09/766,880
 C--> 29 <141> CURRENT FILING DATE:
 W--> 30 2001-01-19
 W--> 35 <150> PRIOR APPLICATION NUMBER:
 W--> 36 US 60/176,839
 W--> 38 <151> PRIOR FILING DATE:
 W--> 39 2000-01-19
 E--> 44 <160> NUMBER OF SEQ ID NOS:
 W--> 45 20
 50 <170> SOFTWARE:
 W--> 51 FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

E--> 56 <210> SEQ ID NO:
 W--> 57 C1
 E--> 59 <211> LENGTH:
 W--> 60 50
 E--> 62 <212> TYPE:
 W--> 63 DNA
 W--> 65 <213> ORGANISM:
 W--> 66 staphylococcus epidermidis
 E--> 71 <400> SEQUENCE:
 72 C1
 73 catgaaccat gtcaggctt gacggaagca gcattaaatg gatcctcata
 74 50
 E--> 78 <210> SEQ ID NO:
 W--> 79 2
 E--> 81 <211> LENGTH:

Rec 1.824 of
 Sequence Header, all
 Sequence Listing files
 MUST be saved in
 Does Not Comply
 Corrected Diskette Needed.
 ASCII
 Text
 format.

The following
 pages show
 samples of
 global errors

FGJ: alphabetical
 leading next
 to numeric
 identifier were
 inserted by CRLF

software

If file is created
 in MS Word, save it
 as MS-DOS Text with Line Breaks

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/766,880A

DATE: 12/13/2002
TIME: 16:12:56

Input Set: A:\PTO.txt
Output Set: N:\CRF4\12132002\I766880A.raw

W--> 82 26
E--> 84 <212> TYPE:
W--> 85 DNA
W--> 87 <213> ORGANISM:
W--> 88 unknown
93 <210> FEATURE:
96 <213> OTHER INFORMATION:
97 *Staphylococcus*
E--> 102 <400> SEQUENCE:
103 2
104 acatgtcagg tcctgacgga agcago
105 26
E--> 109 <210> SEQ ID NO:
W--> 110 3
E--> 112 <211> LENGTH:
W--> 113 50
E--> 115 <212> TYPE:
W--> 116 DNA
W--> 118 <213> ORGANISM:
W--> 119 *Staphylococcus aureus*
E--> 124 <400> SEQUENCE:
125 3
126 catgaaccat gtcaggctct gacggaagca gcattaaatg gatcatata
127 50
E--> 131 <210> SEQ ID NO:
W--> 132 4
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W--> 135 49
E--> 137 <212> TYPE:
W--> 138 DNA
W--> 140 <213> ORGANISM:
W--> 141 unknown
146 <210> FEATURE:
149 <213> OTHER INFORMATION:
150 *Staphylococcus*
E--> 155 <400> SEQUENCE:
156 4
157 catgaaccat gtcaggctct gacggaagca gcattaaatg gatctata
158 49
E--> 162 <210> SEQ ID NO:
W--> 163 5
E--> 165 <211> LENGTH:
W--> 166 46
E--> 168 <212> TYPE:
W--> 169 DNA
W--> 171 <213> ORGANISM:
W--> 172 *streptococcus pyogenes*
E--> 177 <400> SEQUENCE:
178 5

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/766,880A

DATE: 12/13/2002
TIME: 10:12:30

Input Set: A:\PTO.txt
Output Set: N:\CRF4\12132002\I766880A.raw

E--> 274 <400> SEQUENCE:

275 3
276 gtttgtgtg aagggggtaa gggggggaaat ccggcaaccc taaggc
277 46

E--> 281 <210> SEQ ID NO:

W--> 282 10

E--> 284 <211> LENGTH:

W--> 285 41

E--> 287 <212> TYPE:

W--> 288 DNA

W--> 290 <213> ORGANISM:

W--> 291 unknown

295 <220> FEATURE:

296 <223> OTHER INFORMATION:

314 streptococcus

E--> 305 <400> SEQUENCE:

306 10

311 ttgcgtgaag uggtcagggg aggaatccag cagcttaacg

308 41

W--> 313 11

E--> 315 <211> LENGTH:

W--> 316 64

E--> 318 <212> TYPE:

W--> 319 DNA

W--> 321 <213> ORGANISM:

W--> 322 Klebsiella pneumoniae

E--> 327 <400> SEQUENCE:

328 11

329 ccggacgcta ctctgtttac caggcagggt ccggaaaggaa gcggccggcg cagacgacgt

330 ~~gt-gt~~ delete hyphen (-) and insert gt directly beside first group of gt

331 60

332 64

334 64

W--> 340 12

E--> 342 <211> LENGTH:

W--> 343 64

E--> 345 <212> TYPE:

W--> 346 DNA

W--> 348 <213> ORGANISM:

W--> 349 unknown

354 <220> FEATURE:

357 <223> OTHER INFORMATION:

358 ~~Styphimur W~~

E--> 363 <400> SEQUENCE:

364 12

365 ccggacgcta ctctgtttac caggcagggt ccggaaaggaa gcggccggcg cagatgacgc

366 gtgt

367 60

368 64

60

b4

4
RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/766,880A

DATE: 12/13/2002

TIME: 10:12:56

Input Set : A:\PTO.txt
Output Set : N:\CRF4\12132002\I766880A.raw

W--> 569 Artificial Sequence

574 <220> FEATURE:

577 <223> OTHER INFORMATION:

578 ~~oligonucleotide probe~~

E--> 583 <400> SEQUENCE:

584 20

585 tccggggcct tgcataaagtg agtccaggcc ttcatatgttc g

586 41

589 5/5

592 BOSTON 1546047v1

delete